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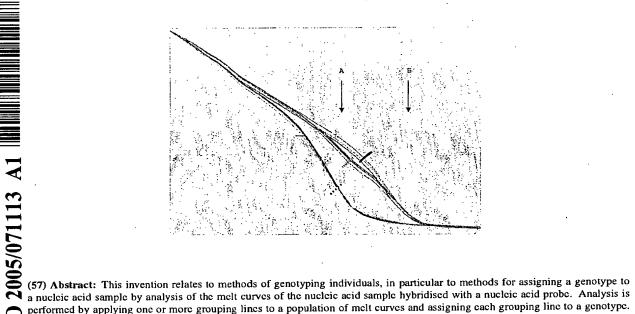
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performed by applying one or more grouping lines to a population of melt curves and assigning each grouping line to a genotype. The genotype of a nucleic acid sample in the population is determined by identifying the grouping line which intersects the melt curve of the nucleic acid sample.

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